

202g	UACC-1012	UACC-1529	UACC-647	WM1791-C	UACC-827	HA-A	UACC-930	UACC-903	UACC-1097	M93-47	TD-1720	TD-1638	TD-1730	solute carrier family 16 (mono carboxylic acid tr
														homeo box D3
														cerebellin 1 precursor
														chemokine (C-X-C motif), receptor 4 (fusin)
														multispanning membrane protein (70kD)
														ataxia telangiectasia and Rad3 related
														profilin 1
														forkhead (Drosophila)-like 1
														ESTs, Highly similar to CGI-03 protein (H sapie
														Homo sapiens mRNA: cDNA DKFZp564B0769
200														ESTs
														ESTs
														ESTs
														RAB5A, member RAS oncogene family
														uno-51 (C elegans)-like kinase 1
														protein tyrosine phosphatase, receptor type, ga
														disabled (Drosophila) homolog 2 (mitogen-resp
														ESTs
														RAB111A, member RAS oncogene family
														antiquitin 1
202r														major histocompatibility complex, class II, Dp b
														fucosyltransferase 8 (alpha (1,6) fucosyltransfe
														tumor protein D52
														cleavage stimulation factor, 3' pre-RNA, subuni

FIG. 1

FIG. 2

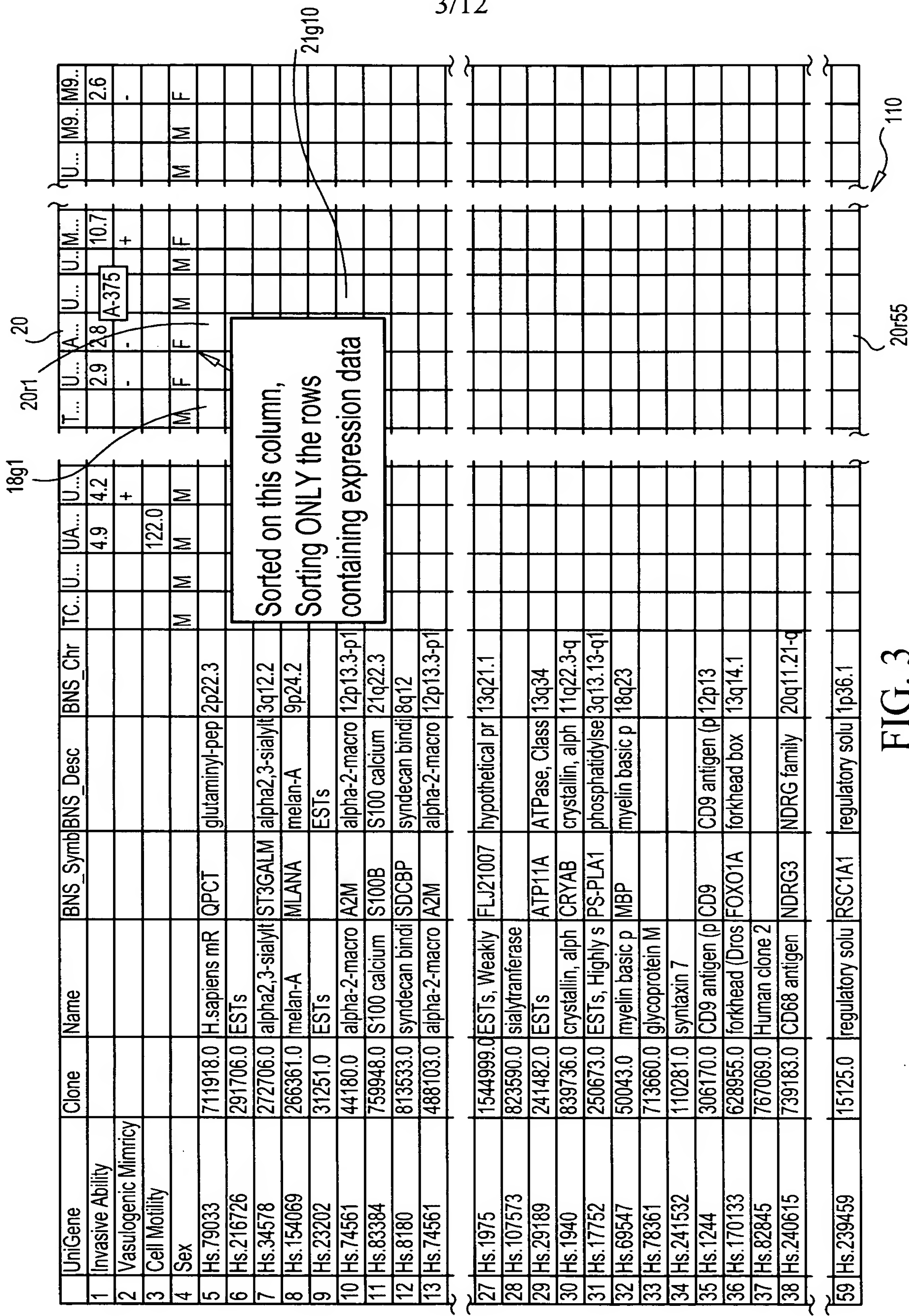


FIG. 3

FIG. 4

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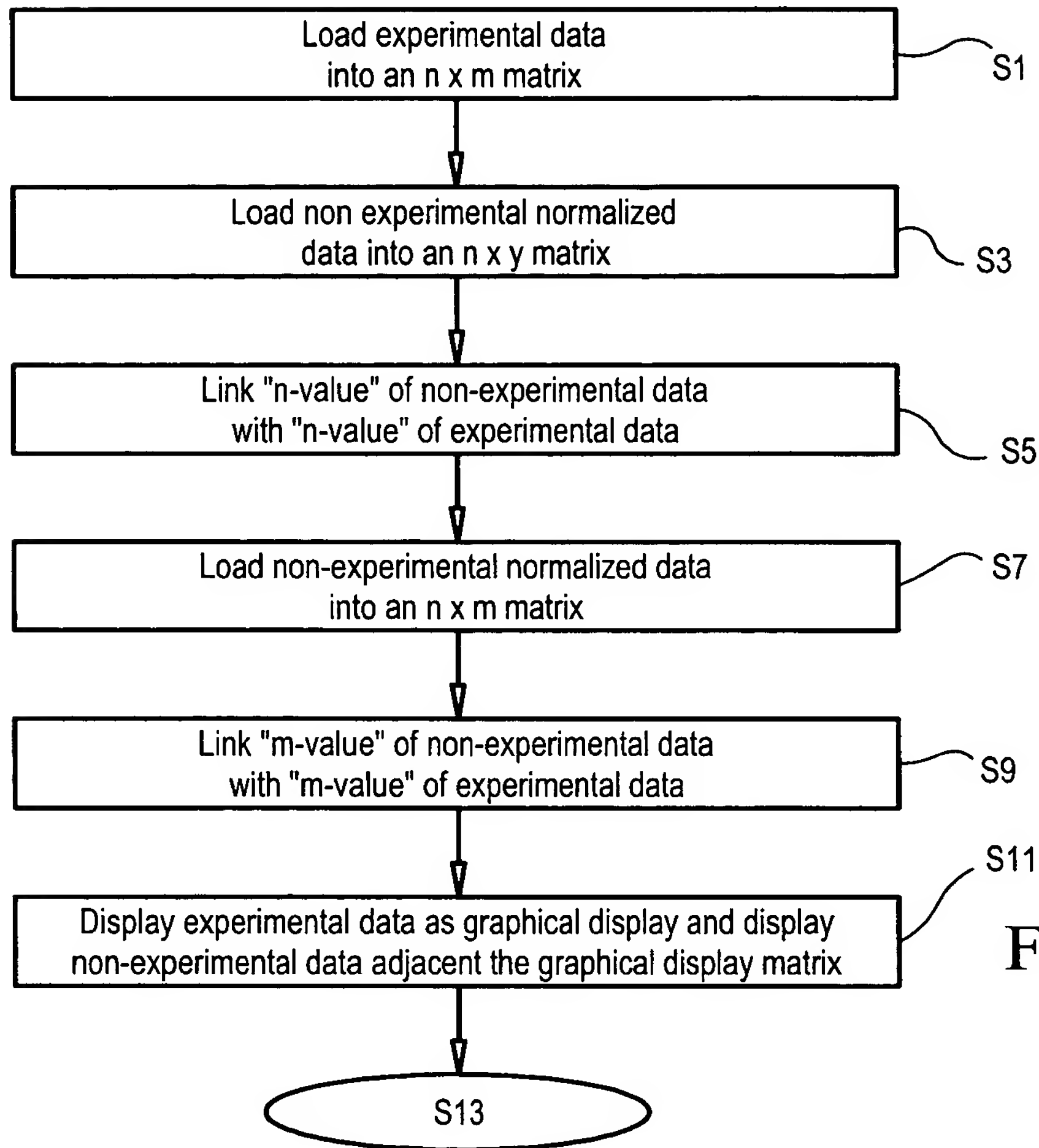


FIG. 5A

	101	102	103
201	101,201	102,201	103,201
202	101,202	102,202	103,202
203	101,203	102,203	103,203
204	101,204	102,204	103,204

FIG. 6A

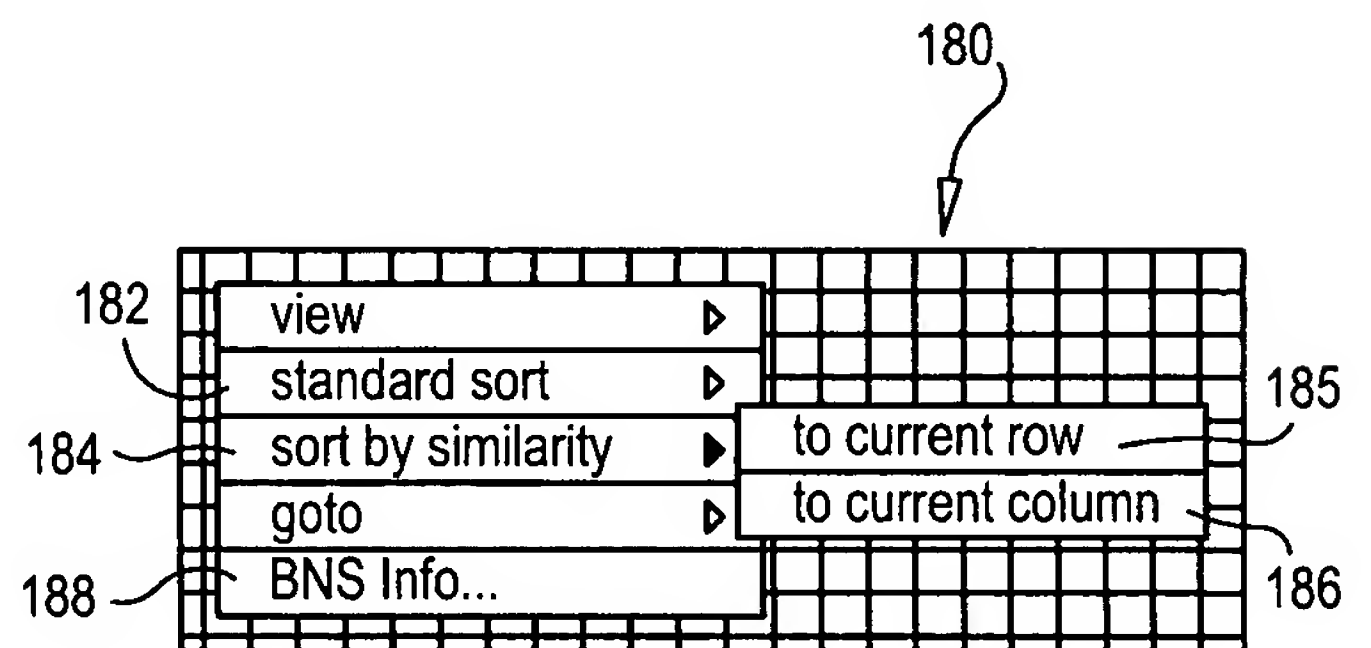
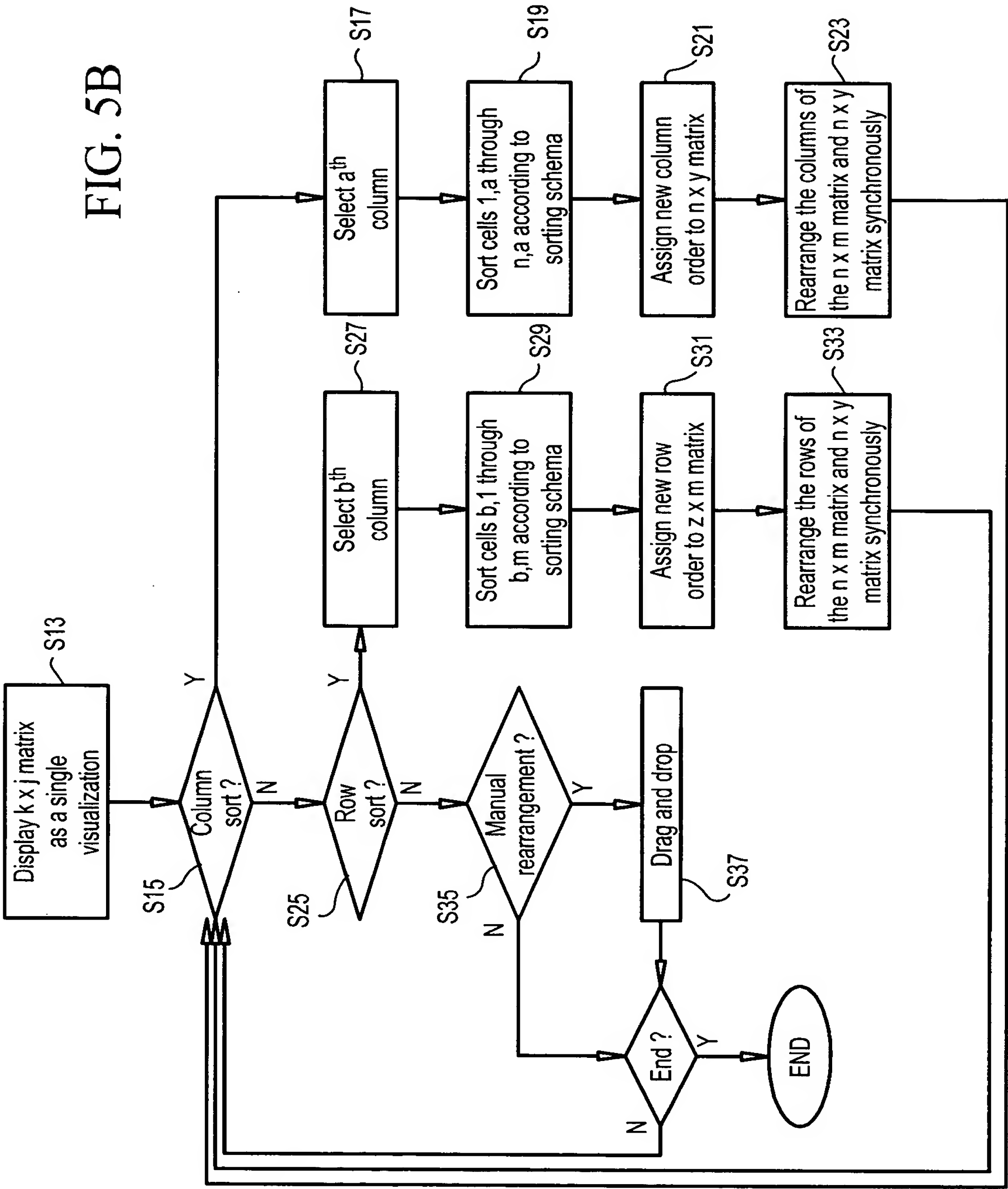


FIG. 6B

FIG. 5B



	101	102	103
202	101,202	102,202	103,202
201	101,201	102,201	103,201
203	101,203	102,203	103,203
204	101,203	102,203	103,203

FIG. 6C

	101	102	103
202	101,202	102,202	103,202
203	101,203	102,203	103,203
201	101,201	102,201	103,201
204	101,204	102,204	103,204

FIG. 6D

Sort Settings

Positive Value

8.0302

(fold increase)

300

Negative Value

8.0304

(fold decrease)

Window Size

2306

(num columns)

Increment

1308

(num columns)

Similarity Increment

100310

(num rows)

OK

Cancel

FIG. 9A

+	+	-		+	-	-	+	+	+	320
8.0	8.0	1/8	1	8.0	1/8	1/8	8.0	8.0	8.0	322
2.1	0.3	0.1	0.2	2.3	0.4	0.1	1.1	0.7	2.2	324
0.3	2.4	0.6	0.1	1.3	0.2	0.2	0.4	1.2	1.0	326
0.3	1.2	0.4	0.1	1.1	0.5	0.2	0.6	1.1	0.8	328

FIG. 9B

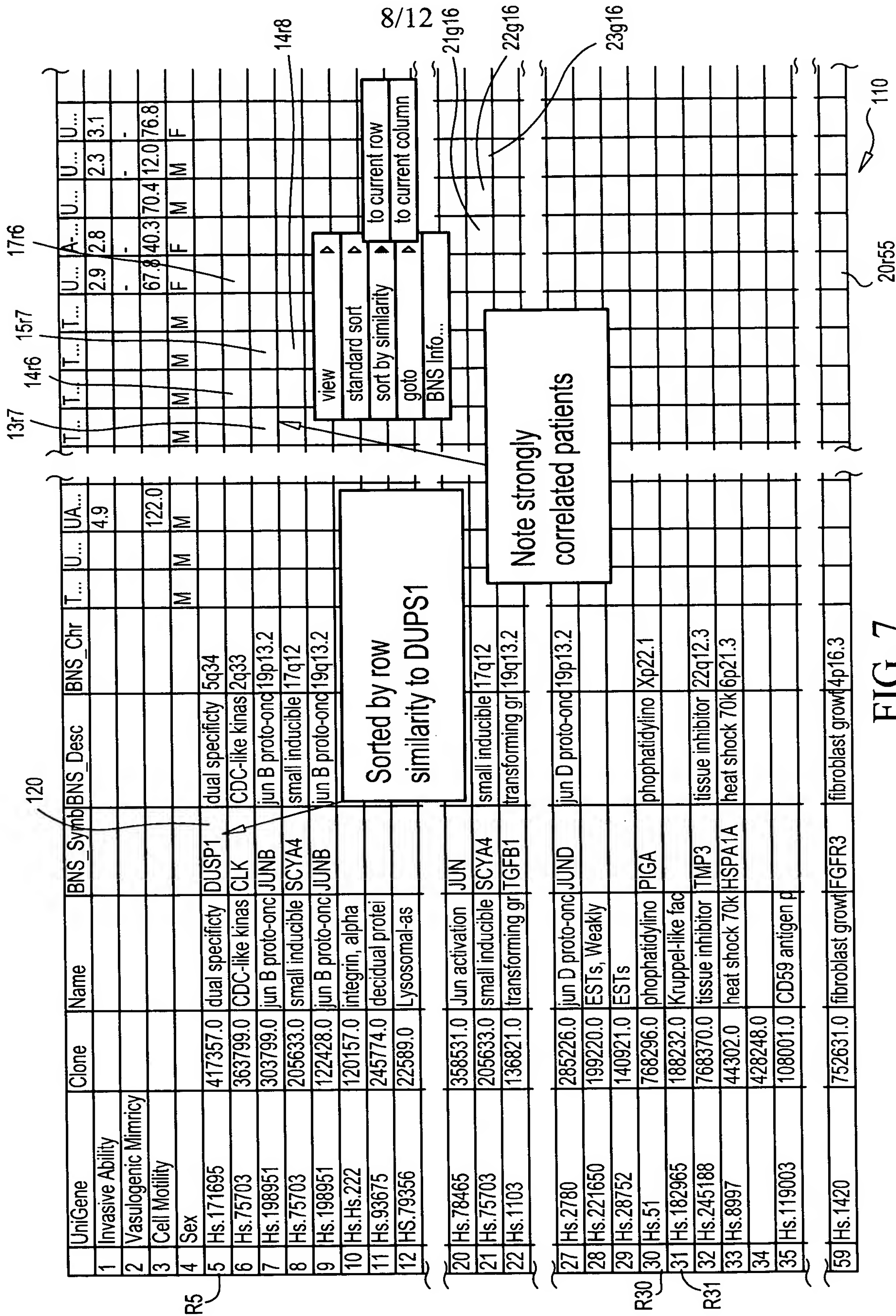


FIG. 7



R6										100										110																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																			
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UniGene	Clone	Name	Symbol	Chr	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g	g

FIG. 10

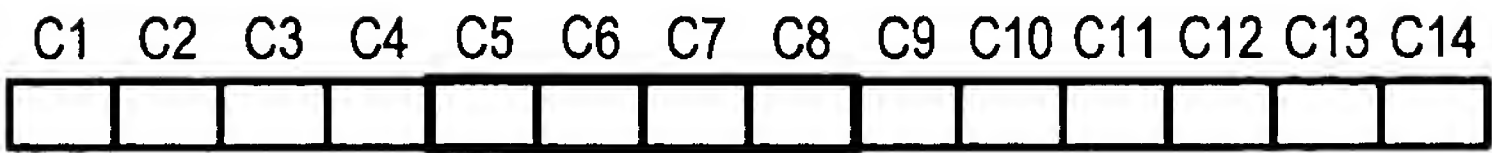


FIG. 11A

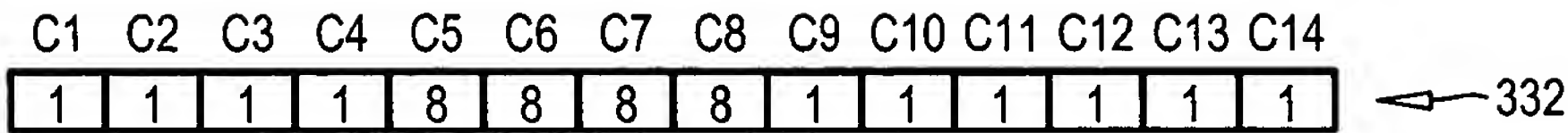


FIG. 11B

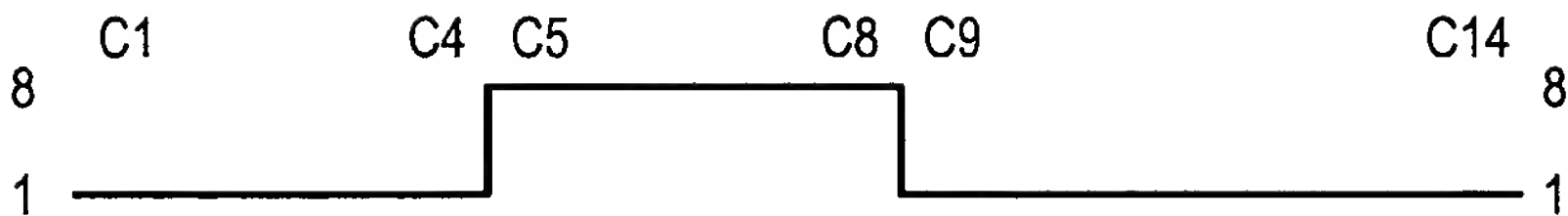


FIG. 11C

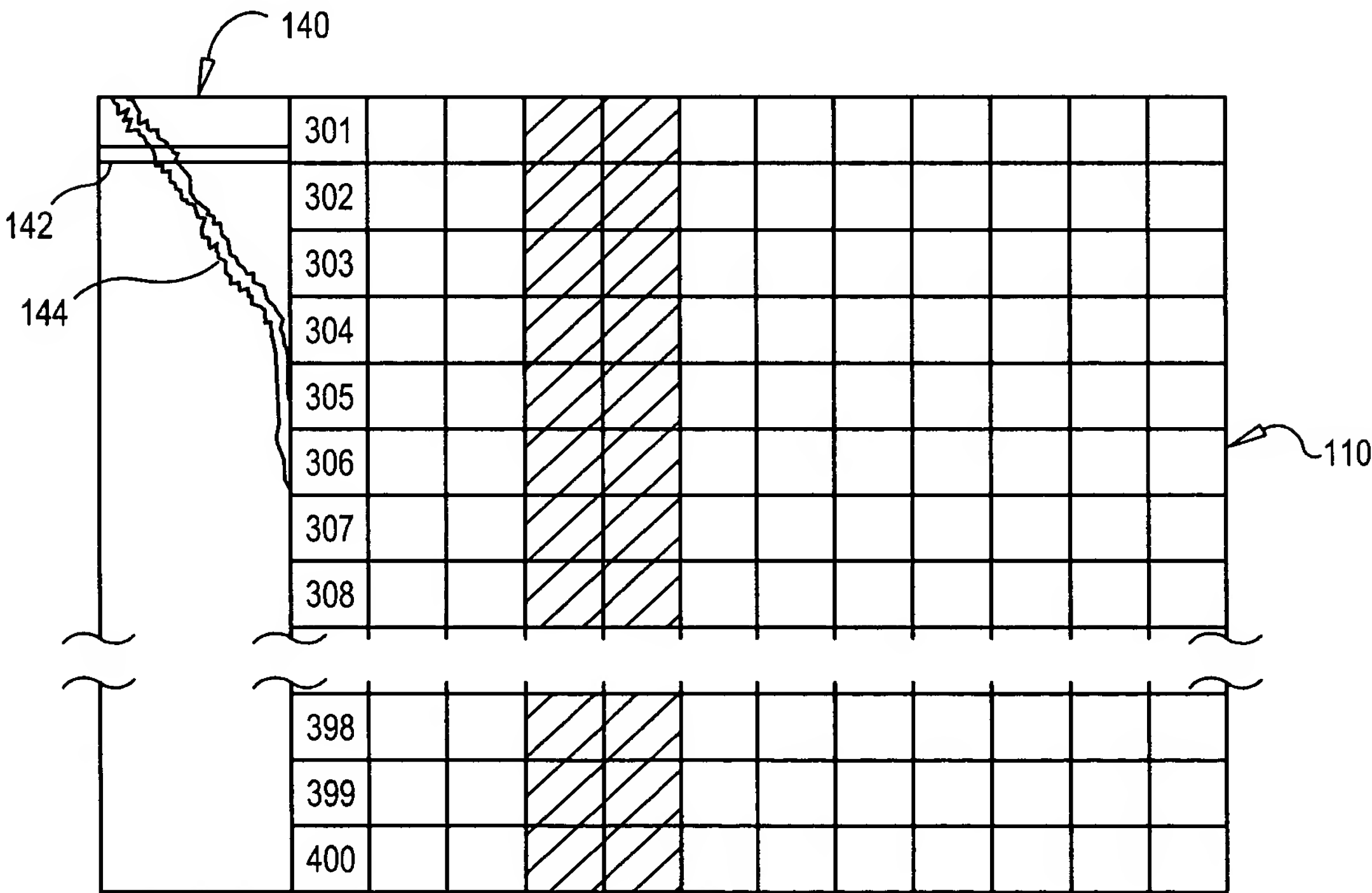


FIG. 13

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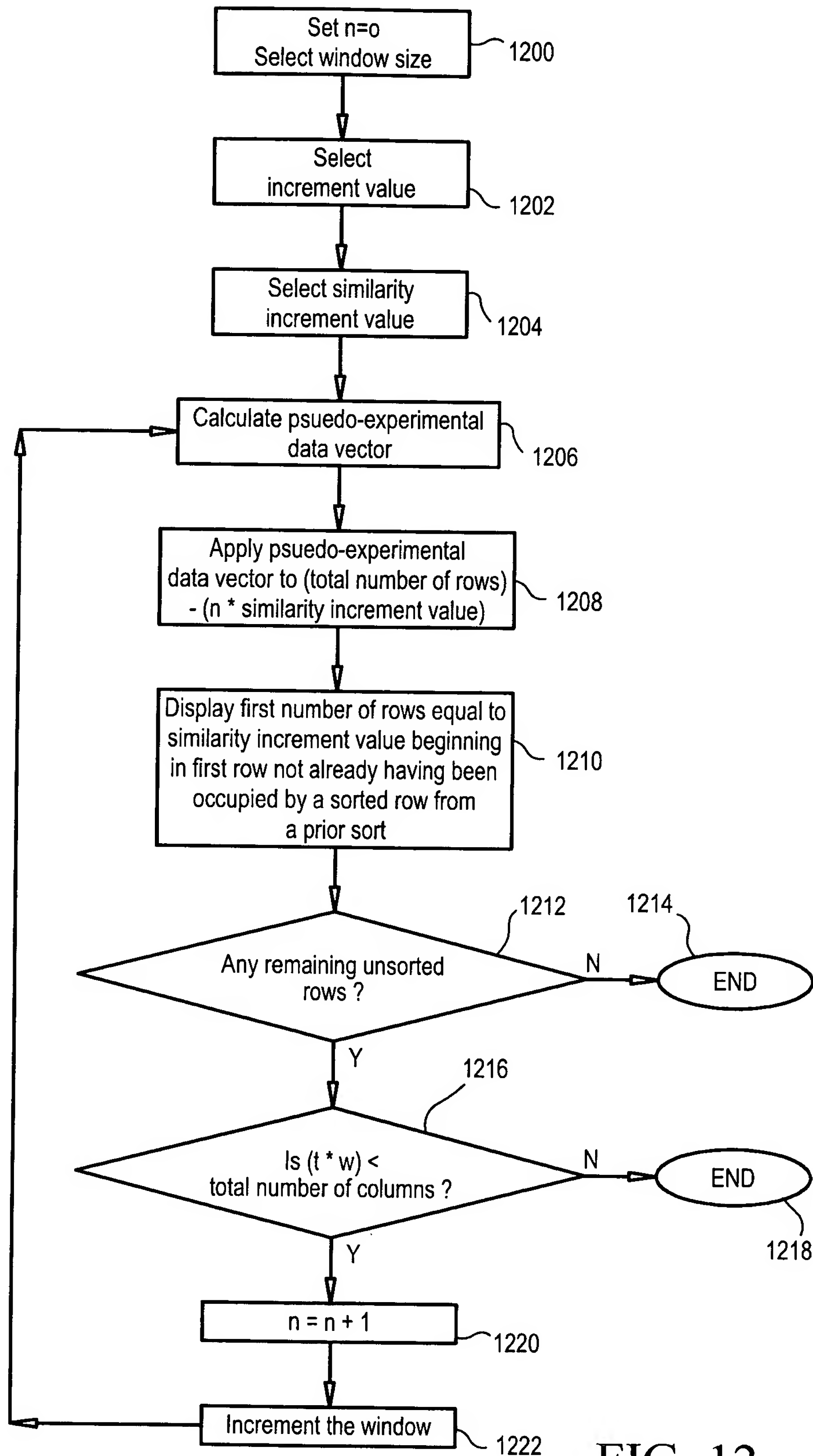


FIG. 12